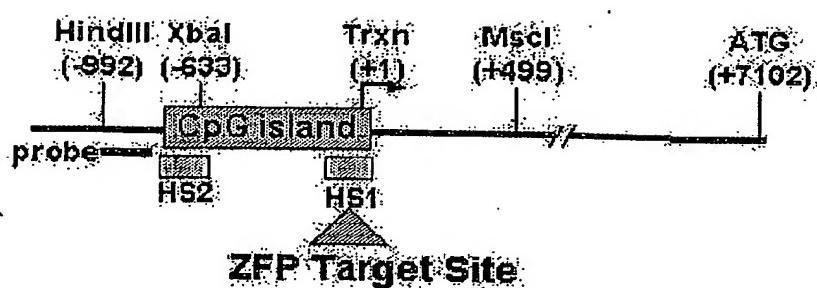
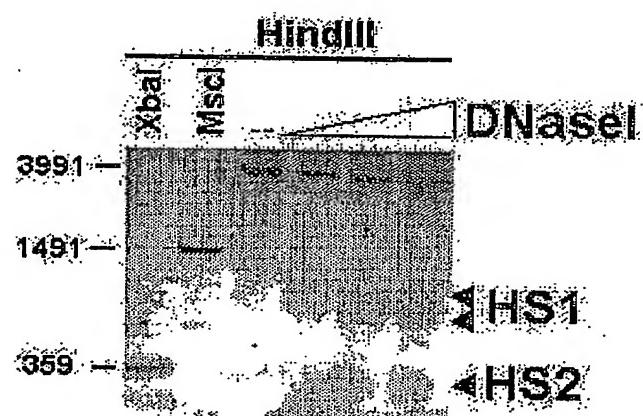
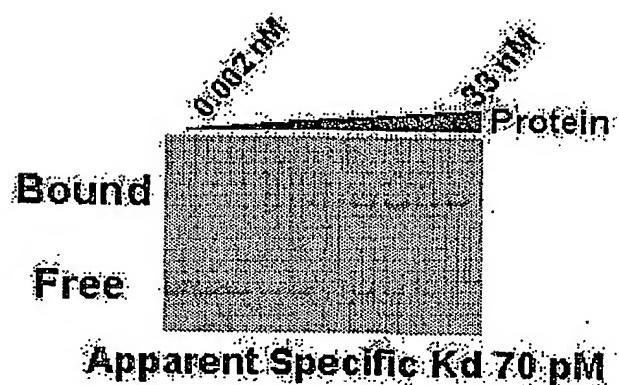


1/15

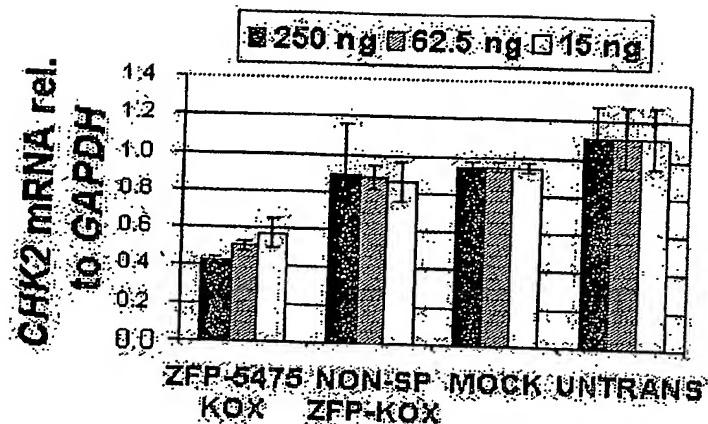
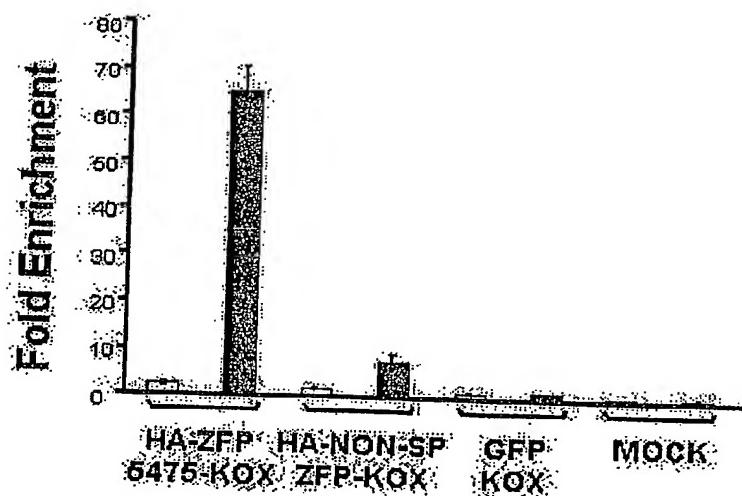
Fig. 1

A**B****C**

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Fig. 2

A**B**

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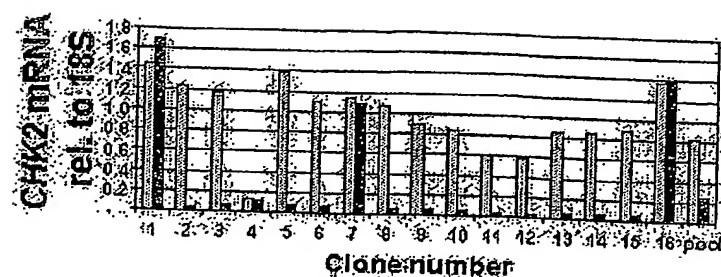
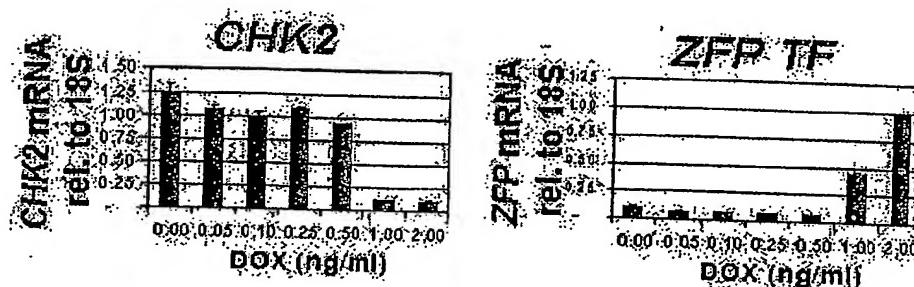
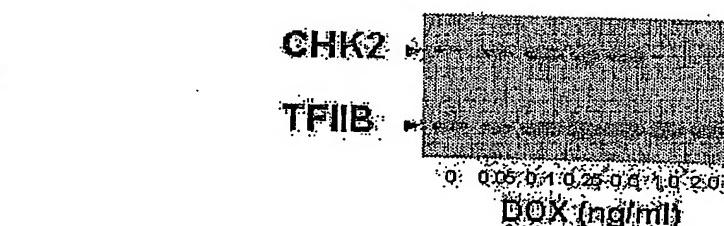
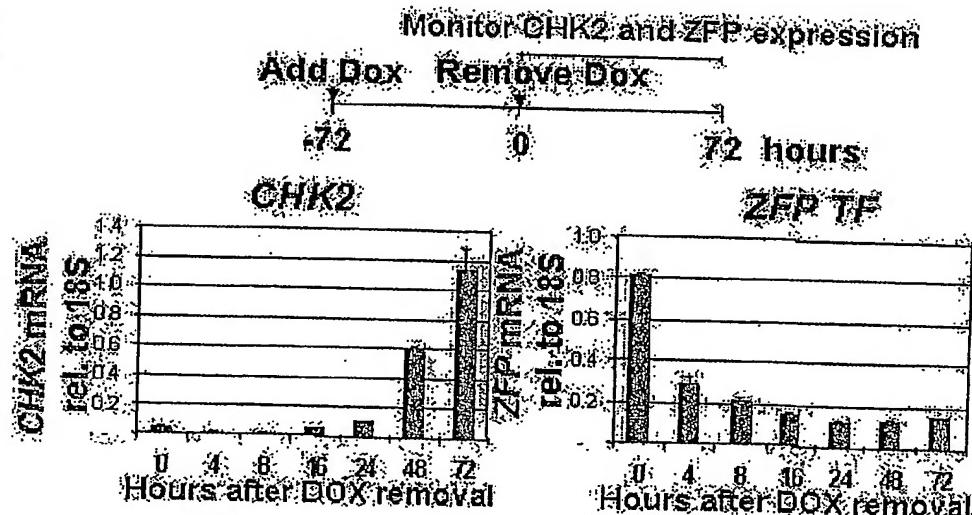
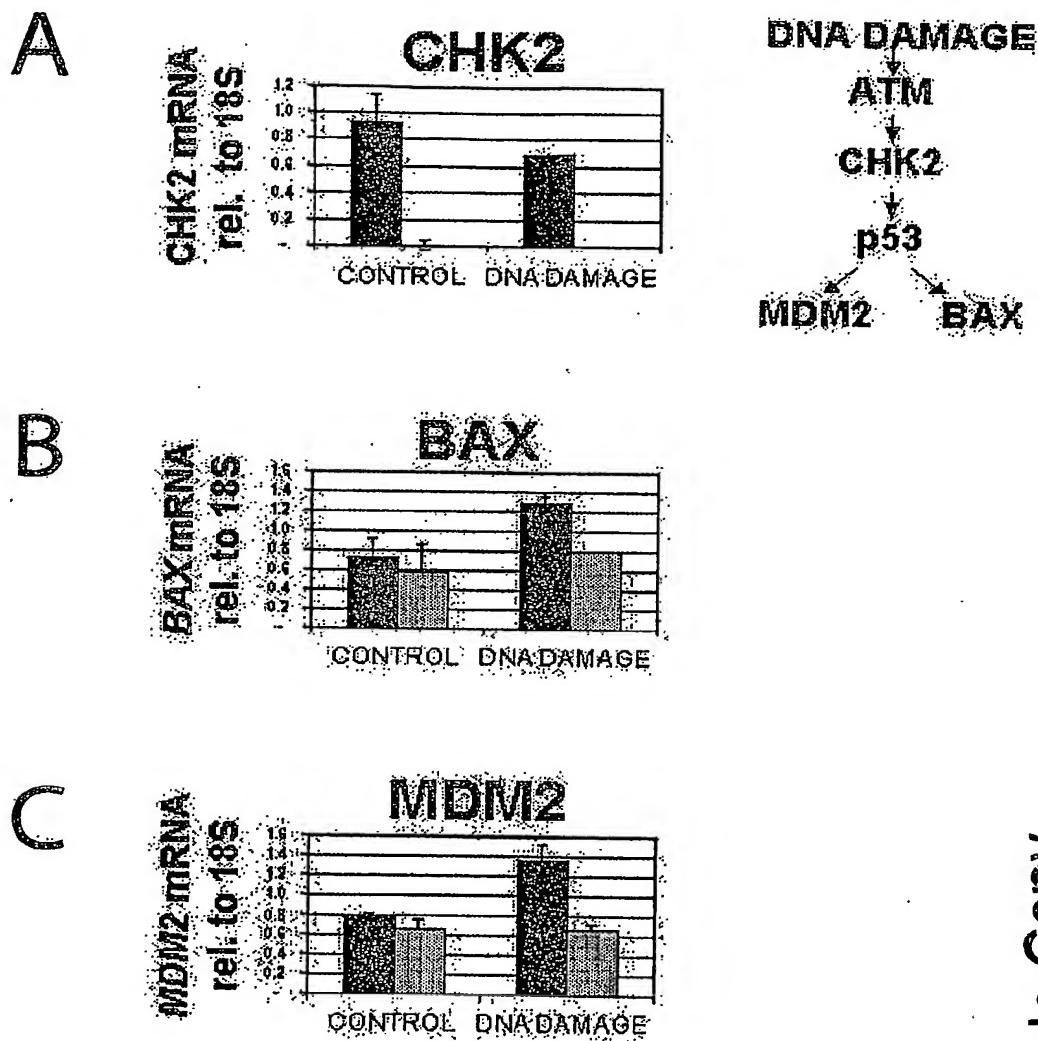
A**B****C****D**

Fig. 3

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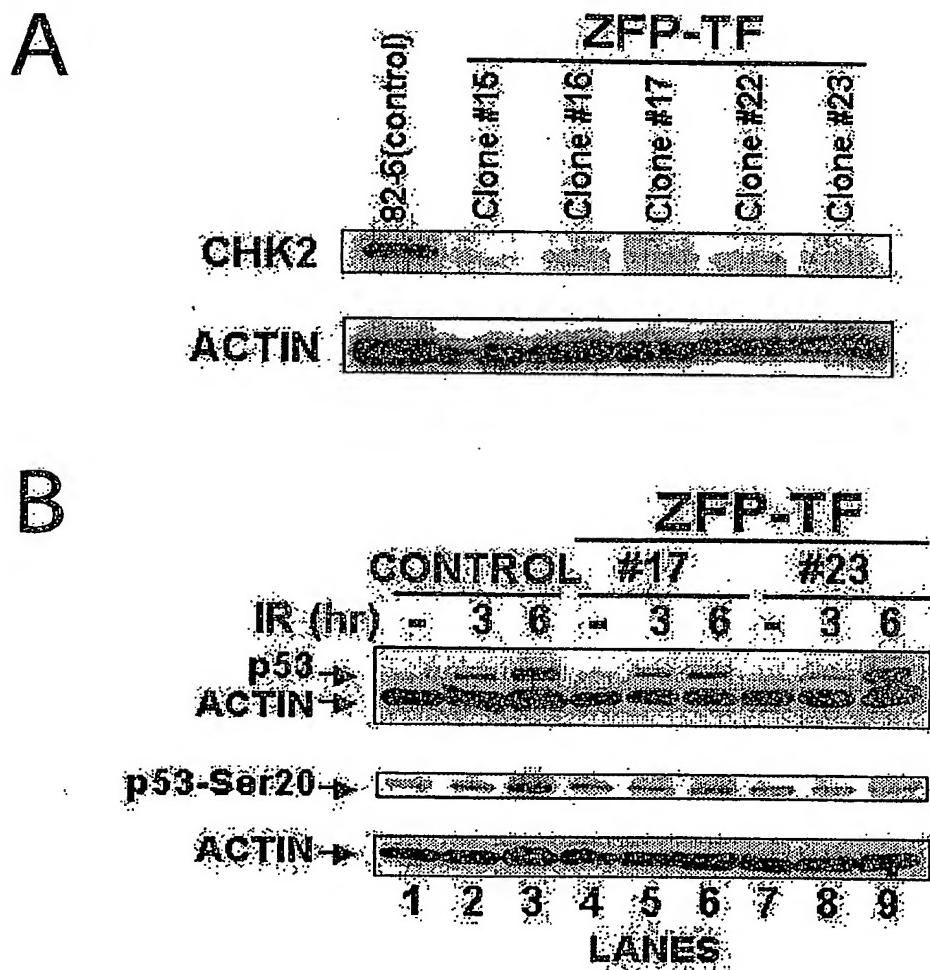
Fig. 4



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Fig. 5



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FIGURE 6

MAERPFQCRICMRNFSRSDHLSRHIRHTGEKPFACDICGRKFADNRDRTKHT
KIHTGGQRPYACPVESCDRRFSDRKTLEHIRIHTGQKPFQCRICMRNFSTSSG
LSRHIRTHTGSQKPFQCRICMRNFSRSDHLSEHIRHTGEKPFACDICGRKFAT
SSDRTKHTKIHLRQKDAARN

SEQ ID NO: 27

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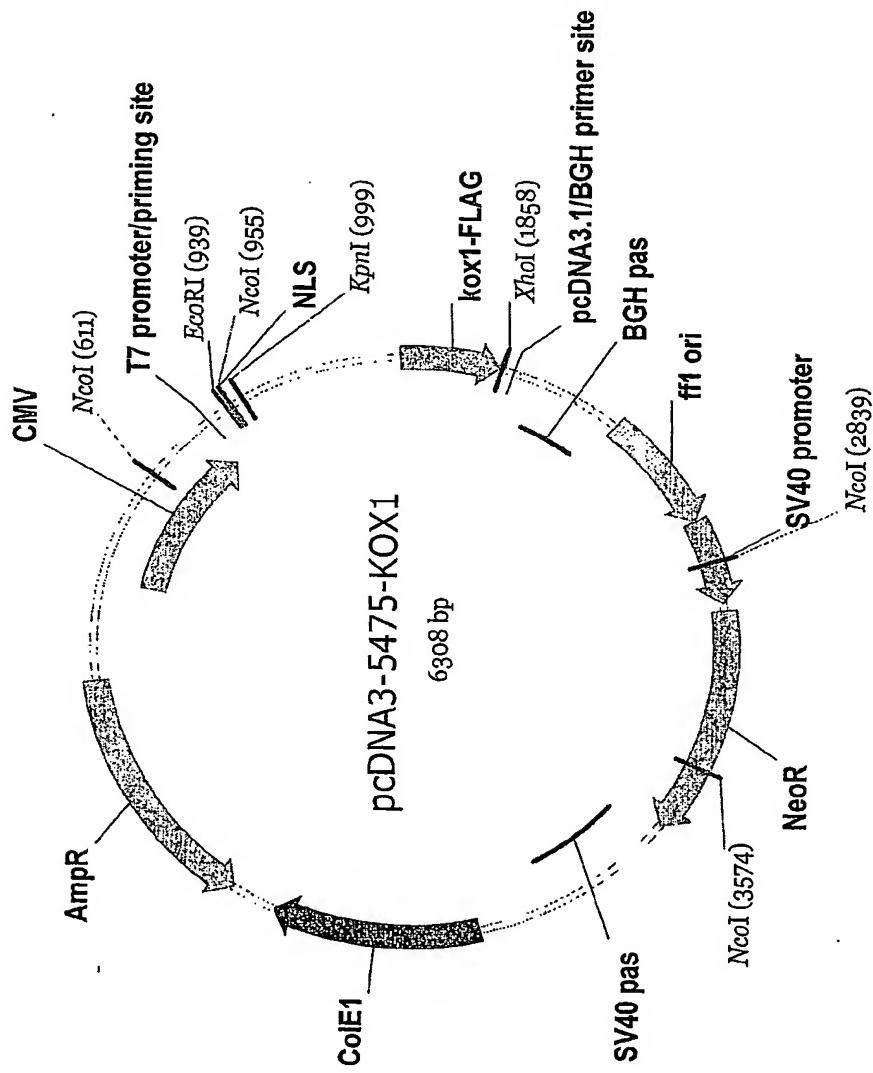
FIGURE 7

MAERPYACPVESCDRRFSTSSADLTEHIRIHTGQKPFQCRICMRNFSAANLSRHIRTHTGGERPF
QCRICMRNFSRSDALSTHIRTHTGEKPFACDICGRKFADRSTRTKHTKIHTGSQKPFQCRICMRN
FSRSDVLSAHIRTHTGEKPFACDICGKKFADRSNRIKHTKIHLRQKDAAR

(SEQ ID NO: 53)

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FIG. 8



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FIG.9A

1 GACGGATCGG GAGATCTCCC GATCCCCAT TGCTGACTCT CAGTACAATC TGCTCTGATG CGGCATAGTT
 CTGCTAGCC CTCTAGAGGG CTAGGGATA CCAGCTGAGA GTCATGTTAG ACAGAACTAC GGCGTATCAA
 71 AAGCAGTAT CTGCTCCCTG CTGTTGTGT GGAGGTGGCT GAGTAGTGC CGAGCAAAAT TTAAGGTCTACA
 TTTCGGTCATA GACGAGGGAC GAACACACAA CCTCCAGCGA CTCATCACGC GCTCGTTTA ATTICGATGT
 141 ACAAGGCAAG GCTTGACCGA CAATTGCATG AAGAATCTGC TTAGGGTTAG GCCTTTCGCG CTGCTTCGCG
 TGTTCGGTC CGAACCTGGCT GTTAACGTAC TTCTTAGACG AATCCAATC CGAAAACGC GACGAAGCGC
 211 ATGTAACGGGC CAGATATAGC CGTTGACATT GATTATGAC TAGTATATAA TAGTAATCAA TTACGGGTCTC
 TACATGCCCG GTCTATATGC GCAACTGTAA CTAATAACTG ATCAATAATT ATCATATTAGT ATGCCCCAG
 281 ATTAGTTCAT AGCCCATATA TGGAGTTCCG CGTTACATAA CTTACGGTAA ATGGCCGCC TGGCTGACCG
 TAATCAAGTA TCGGGTATATA ACCTCAAGGC GCAATGTATT GAATGCCATT TACCGGGGG ACCGACTGGC
 351 CCCAACGACC CCCGCCATT GACGTCATAA ATGACGTATG TTCCCATAGT AACGCCAATA GGGACTTCC
 GGGTGTGG GGGGGGAA CTGCAAGTT TACTGCATAC AAGGGTATCA TTGCGGTAT CCGTGAAGG
 421 ATTGACGTCA ATGGGGAC TATTACGGT AAACGTCCCA CTTGGCAGTA CATCAAGTGT ATCATATGCC
 TAACTGGCAGT TACCCACCTG ATAATGCCA TTGACGGGT GAACGTCAAT GTAGTTCACCA TAGTATACGG
 491 AAGTACGCC CCTATTGACG TCAATGACGG TAAATGCCC GCCTGGCATT ATGCCAGTA CATGACCTTA
 TTCATGCGGG GGATAACTGGC AGTTACTGCC ATTACCGGG CGGACCGTAA TACGGGTAT GTACTGGAT

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561 TGGGACTTTC CTACTTGGCA GTACATCTAC GTATTAGTC CTCGCTATTAC TCGCTATTAC CATGGTGTGATG CGGTGTTGGC
 ACCCTGAAAG GATGAAACGGT CATGTTAGATG CATAATCAGT AGCGATAATG GTACCACTAC GCCAAACCG
 631 AGTACATCAA TGGGGGTGCA TAGGGTTG ACTCACGGGG ATTICCAAGT CTCACCCCA TTGACGTCAA
 TCATGTAGTT ACCGGCACT ATCGGCCAAC TGAGTGCCTC TAAAGGTCTCA AAATGTCGTA ACAACTCCGC CCCATTGACG
 701 TGGGAGTTG TTTGGCAAC AAAATCAACG GAGACTTCAC TTTTAAAGGTGC CTCGAAAGT TTTACAGCAT TGTGAGGGC GGGTAACTGC
 ACCCTCAAAAC AAAACCGTGG TTTAGTTGC CTCGAAAGT GCTCTATCAA GCAGAGCTCT CTTGGCTAACT AGAGAACCCA
 771 CAAATGGGG GTAGGGGTG ACGGTGGGAG GCTCTATCAA CGAGGAGCTCT CTTGGCTAACT AGAGAACCCA
 GTTITACCGC CATCCGCACA TGCCACCCCTC CAGATATTT CGTCTCGAGA GACCGATTGA TCTCTGGGT
 841 CTGCTTACTG GCTTATCGAA ATTAAATACGA CTCACATAG GGAGACCCAA GCTGGCTAGC GTTAAACTT
 GACGAATGAC CGAATAGCTT TAATTATGCT GAGTGTATTC CCTCTGGGT CGACCGATCG CAAATTGAA

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ECORI ~~~~~

M A P K K R K V .

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FIG.9B

911 AAGCTGATCC ACTAGTCAG TGTGGTGGAA TTCGCTAGCG CCACCATGGC CCCAAGAA AGAGGAAGG
TTCGACTAGG TGATCAGTC ACACCACCTT AAGCGATCGC GGTGGTACCG GGGTTCTTC TTCTCCCTTC

KpnI

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981 TGGGAATCGA TGGGGTACCC TTCCAGTGTG GAATCTGCAT GCGTAACCTTC AGTCGTAGTG ACCACCTGAG  
ACCCCTTAGCT ACCCCATGGG AAGGTACAG CTTAGACGTA CGCATTTGAAG TCAGCATCAC TGGTGGACTC  
R H I R T H T G E K P F A C D I C G R K F A D

1051 CCGGCACATC CGCACCCACA CAGGGAGAA GCCTTTTIGCC TGTGACATT GTGGAGGAA ATTGGCCGAC  
GGCCGTGTAG GCGTGGGTGT GTCCGGCTCTT CGGAAAACGG ACACTGAAA CACCTCTT TAAACGGCTG  
N R D R T K H T K I H T G G Q R P Y A C P V E S

1121 AACCGGGACC GCACAAAGCA TACCAAGATA CACACGGGG GACAGGGGCC GTACGCATGC CCTGTGCGAGT  
TTGGCCCTGG CGTGTTCGTT ATGGTTCTAT GTGTGCCCG CTGTGCCCG CATGCGTAG GGACAGCTCA  
C D R R F S D R K T L I E H I R I H T G Q K P

1191 CCTGGGATCG CGCGCTTTCT GACAGGAAGA CACTTATCGA GCATATCCGC ATCCACACCG GTCAAGGCC  
GGACGCTAGC GGGGAAAGA CTGTCCTCT GTGAATAGCT CGTATAGGC TAGGTGTGGC CAGTCTTCGG

F Q C R I C M R N F S T S G L S R H I R T H

1261 CTTCCAGTGT CGAACATCGCA TGGCTTAACCT CAGTACCGC AGCGGGCTGA GCCGCCACAT CGGCCAC  
GAAGGGTACA GCTTAGACGT ACGGATTGAA GTCATGGTCG TCGCCCGACT CGGGGGTGA GGGTGGGTG

T G S Q K P F Q C R I C M R N F S R S D H L S E

1331 ACAGGATCTC AGAAGGCCCTT CCAGTGTGCGA ATCTGCATGC GTAACTTCAG TCGTAGTGAAC CACCTGAGCG  
TGTCTAGAG TCTTCGGAA GGTCAACAGCT TAGACGTAAG CATTGAAGTC AGCATCACTG GTGGACTCGC  
H I R T H T G E K P F A C D I C G R K F A T S

1401 AACACATTG CACCCACACA GGGGAGAAC CTTTGCCTG TGACATTGT GGGAGGAAT TTGCCACACAG  
TTGTGTAAGC GTGGGGTGTG CCGCCTCTCG GAAACGGAC ACTGTAACAA CCCTCCCTTA AACGGGGTC  
S D R T K H T K I H L R Q K D A A R G S G M D

1471 CAGGGACCGC ACAAGGCATA CCAAGATACA OCTGGCCCAA AAAGATGCGG CCCGGGGATC CGGCATGGAT  
GTCGCTGGCG TGTTTCGTAT GGTTCATGT GCACTGGGTT TTTCTACGCC GGGCCCCTAG GCCGTACCTA  
A K S L T A W S R T L V T F K D V F V D F T R E

1541 GCTAAAGTCAC TAATGCGCTG GTCCCCGACA CTGGTGACTC TCAAGGATGT ATTGTGGAC TTCACCAAGGG  
CGATTCACTG ATGACGGAC CAGGGCTGT GACCACTGGA AGTACCTACA TAAACACCTG AAGGGTCCC

## FIG.9C

1611 E W K L L D T A Q Q I V Y R N V M L E N Y K N  
 AGGAGTGGAA GCTGCTGGAC ACTGCTCAGC AGATCGTGT CAGAAATGTG ATGCTGGAGA ACTATAAGAA  
 TCCCTACCTT CGACGACCTG TGACGACTG TCTAGCACAT GTCTTACAC TACGACCTTG TGATATTCTT  
 L V S L G Y Q L T K P D V I L R I E K G E E P  
 CCTGGTTTC TTGGGTTATC AGCTTAAGTAA GCCAGATGTG ATCCCTCCGGT TGGAGAAGGG AGAAGAGCCC  
 GGACCAAAGG AACCCAAATAG TCGATGATT CGGTCTACAC TAGGAGGCCA ACCTCTTCCC TCTTCTCGGG  
 W L V E R E I H Q E T H P D S E T A F E I K S S  
 1751 TGGCTGGTGG AGAGAGAAAT TCACCAAGAG ACCCATCTG ATTCAAGAGAC TGCATTGGAA ATCAAATCAT  
 ACCGACCACC TCTCTCTTA AGTGGTTCTC TGGTAGGAC TAAGTCTG ACGTAAACTT TAGTTAGTA

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XbaI

1821 CAGTGACTA CAAGGACGAC GATGACAAGT AAGCTTCTCG AGTCATGCTA GAGGGCCGT TTAAACCCGC
 GTCAACTGAT GTTCCTGCTG CTACTGTTCA TTGGAAGAGC TCAGATCGAT CTCCCCGGCA AATTGGCG
 1891 TGATCAGCCT CGACTGTGCC TTCTAGTGTG CAGCCATCTG TTGTTGCGCC CTTCTCTTGA
 ACTAGTCGGA GGTGACAGG AAGATCAACG GTGGGTAGAC AAACAACGGG GAGGGGGCAC GGAAGGAACCT
 1961 CCCTGGAAAG TGGCACTCCC ACTGTCTTT CCTAATAAA TGAGGAAATT GCATCGCATT GTCTGAGTAG
 GGGACCTTCC ACGGTGAGGG TGACAGAAA GGATTATTTC ACTCTTTAA CGTAGCTTAA CAGACTCATC
 2031 GTGTCAATTCT ATTCTGGGG GTGGGGTGGG GCAGGACAGC AAGGGGGAGG ATTGGGAAAGA CAATAGCAGG
 CACAGTAAGA TAAGCCCC CACCCCAACCC CGTCCCTGTG TTCCCCCTTC TAACCCCTTCT GTTATCGTCC
 2101 CATGCTGGG ATGCGGTGGG CTCTATGGCT TCTGAGGGG AAAGAACAG CTGGGGCTCT AGGGGGTATC
 GTACGACCCC TAGGCCACCC GAGATACCGA AGACTCCGCC TTCTCTGGTC GACCCCGAGA TCCCCCATAG
 2171 CCCACGCGCC CTGTAGCGGC GCATTAAGCG CGTAAATTGCG GGGGGGTGTG GGTGGTTACG CGCGTACACT
 GGGTGGCGG GACATGCCG GCGCCCCACA CCACCAATGC GCGTGGCACT GGCATGTGA
 2241 TGCCAGGCC CTAGGCCCG CTCCCTTCGC TTCTCTCCCT CCCTTCTCG CACGTTCTCG CGGTTTCCC
 ACGGTGGCGG GATCGGGGC GAGGAAAGCG AAAGAAGGA AGGAAGAGC GTGGCAAGGC CGCGAAAGGG
 2311 CGTCAAGCTC TAAATGGGG CATCCCTTA GGGTTCCGAT TTAGTGCCTT ACGGCACCTC GACCCAAAA
 GCAGTCCGAG ATTAGCCCC GTAGGGAAAT CCCAAGGCTA AATCACGAAA TGCCTGGAG CTGGGGTTT
 2381 AACTIGATTA GGGTGTATGGT TCACGTAGTG GGCCATCGCC CTGATAGACG GACTATCTGC CAAAAGGG GAAACTGCAA
 TTGAACTAAT CCCACTACCA AGTGCATCAC CGGTAGGG GAAACAAACAC TCAACCCTAT CTCGGTCTAT

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FIG. 9D

CCTCAGGTGC AAGAAATTAT CACCTGAGAA CAAGGTTGA CCTTGGCTG AGTTGGATA GAGCCAGATA
 2521 TCTTTGATT TATAAGGAT TTGGGGATT TCGGCCTATT GGTTAAAAA TGAGCTGATT TAACAAAAAT
 AGAAAACCAA ATATTCCTTA AAACCCCTAA AGCCGGATAA CCAATTCTT ACTCGACTAA ATTGTTTA
 2591 TTAACGCAGA TTAATTCTGT GGAATGTGTG TCAGTGTAGG TGTGAAAGT CCCAGGGTC CCCAGGCAGG
 AATTGCGCTT ATTAAGACA CCTTACACAC AGTCAATCCC ACACCTTCA GGGTCCGAG GGGTCCGTCC
 2661 CAGAAGTATG CAAAGCATGC ATCTCAATT GTCAACC ACCAGAACC AGTGTGGAA AGTCCCCAGG CTCCCCAGCA
 GTCTCTATAC GTTTCGTCAG TAGAGTTAAT CAGTCGTGG TCCACACCTT TCAGGGTCC GAGGGTGTG
 2731 GGCAGAAAGTA TGCAAAGCAT GCATCTCAAT TAGTCAGCAA CCATAGTCCC GCCCCTAACT CGGCCCATCC
 CCGTCTTCAT ACGTTTCGTA CGTAGAGTTA ATCAGTCGTT GGTATCAGGG CGGGGATTGA GGCGGTAGG

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2801 CGCCCTAAC TCCGCCCTAGT TCCGCCATT CTCCGCCATT TGGCTGACTA ATTTCCTTAA TTATGCGA  
 GCGGGGATTG AGGGGGTCA AGGGGGTAA GAGGGGGGT ACCGACTGAT TAAAAAAAT AAATACGTCT  
 2871 GCGCGAGGCC GCGCTCTGGCT CTGAGCTATT CCAGAAAGTAG TGAGGGGCT TTTTGGAGG CCTAGGCTTT  
 CCGGCTCCGG CGGAGACCGA GACTCGATAA GGTCTTCATC ACTCCCTCGA AAAACCTTC GGATCCGAAA  
 2941 TGCAAAAGC TCCCGGGAGC TTGTATATCC ATTTCGGAT CTGATCAAGA GACAGGGATGA GGATCGTTTC  
 ACGTTTTTCG AGGGCCCTCG AACATATAGG TAAAGGCCAA GACTAGTCTT CTGTCCTACT CCTAGCAAAG  
 3011 GCATGATTGA ACAAGATGGA TTGCAACCGAG GTTCTCCGGC CGCTTGGGT GAGAGGTAT TCGGGCTATGA  
 CGTACTAACT TGTCTTACT AACGTGCGTC CAAAGGGCG GCGAACCCAC CTCTCCGATA AGCCGATACT  
 3081 CTGGCACAA CAGACAATCG GTCGCTCTGA TGCCGCCGTG TTCCGGTGT CAGGCCAGGG GCGCCGGTT  
 GACCCGGTT GTCTGGTTAGC CGACCGAGCT ACCGGGGCAC AAGGGGACA GTGCCGGTCC CGGGGCCAA  
 3151 CTTTGTCA AGACCGACCT GTCCGGTGGC CTGAATGAAC TGCAGGACGA GGCAGCGGG CTATCGGGC  
 GAAAACAGT TCTGGCTTGA CAGGCCACGG GACTTACTTG ACGTCTCTGCT CGTCTGGCC GATAGCACCG  
 3221 TGGCACGAC GGGCGTTCTC TGGCAGGTG TGCTCGACGT TGTCACTGAA GGGGAAGGG ACTGGCTGCT  
 ACCGGTGTG CCGCAAGGA ACGGGTGGAC ACGAGGTGCA ACAGTGAATT CGCCCTTCCTC TGACCGACGA  
 3291 ATGGGGAA GTGCCGGGG AGGATCTCCT GTCATCTCAC CTTGCTCCCTG CGAGAAAGT ATCCATCATG  
 TAACCCGGTT CACGGCCCCG TCCTAGAGGA CAGTAGAGTG GAACGAGGAC GGCTCTTICA TAGGTAGTAC  
 3361 GCTGATGCCA TGCGGGGGCT GCATAGCTT GATCCGGCTA CCTGCCATT CGACCAAAA GCGAAZACATC  
 CGACTACGTT AGGCCGCCA CGTAGGCAGA CTAGGCCGAT GGACGGGTAA GCTGGTGGTT CGCTTGTGAG  
 3431 GCATCAGGG AGCACGTTACT CGGATGGAAG CCGGTCTTGT CGATOAGGAT GATCTGGACG AAGAGCATCA  
 CGTAGCTCGC TCGTGCATGA GCTACCTTC GCCTACCTCA CTAGACCTGC TTCTCGTAGT

## FIG.9E

|      |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 |
|------|-----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| 3501 | GGGGCTCGCG CCAGCGAAC TGTCGCCAG GCTCAAGGGG CGCATGCCG ACGGGAGGA TCTCGTGTG<br>CCCCGAGCGC GGTCGGCTG ACAAGGGTC CGAGTCCGC CGTACGGG TGCGCTCCT AGAGGACAC<br>NcoI                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        |
| 3571 | ~~~~~ ACCCATGGCG ATGCCTGCTT GCCGAATAATC ATGGGGAAA ATGGCCGCTT TTCTGGATTC ATCGACTGTG<br>3641 TGGGTACCGC TAGGGACGAA CGGCTTATAG TACCACTTT TACCGGGCAA AAGACCTAAG TAGCTGACAC<br>CGGGCGTGGG TGTGGGGAC CGCTATCAGG ACATAGCGGT GGCTACCGT GATATTGCTG AAGAGCTTGG<br>CGGCCGACCC ACACCGCTG GCGATAGTCC TGTATCGAA CCGATGGCA CTATAACGAC TTCTCGAAC<br>3711 CGGGCAATGG GCTGACCGT TCCCTGIGCT TIACTGGTATC GCGCTCCCG ATTGCGAGG CATGCCCTTC<br>GCCGCTTAC CGACTGGCGA AGGAGCACGA AATGCCATAG CGGGAGGGC TAAGCGTGC GTAGCGGAAG<br>3781 TATGCCCTTC TTGACGAGTT CTTCCTGAGCG GGACTCTGGG GTTCAAAATG ACCGACCAAG CGACGCCAA<br>ATAGGGGAAG AACTGCTAA GAAGACTCGC CCTGAGACCC CAAGCTTAC TGGCTGTTT GCTGCGGGTT<br>3851 CCTGCCATCA CGAGATTTCG ATTCCACCGC CGCCTTCTAT GAAAGGTTGG GCTTCGGAAAT CGTTTCCGG<br>GGACGGTAGT GCTCTAACGC TAAGGGGGC GCGGAAGATA CTTTCACACC CGAACGGCTTA GCACAAAGGCC<br>3921 GACGGGGCT CGATGATCTT CCAGGGGG GATCTCATGC TGGAGTCTT CGGCCACCCC AACTGTTA<br>CTGGGGCGGA CCTACTAGGA GGTGGGGCC CTAGAGTACG ACCTCAAGAA GCGGGTGGG TTGACAAT<br>3991 TTGCAAGCTTA TAATGGTTAC AAATAAAGCA ATAGCATAC AAATTCAAAATAAGCAT TTTTTCACT<br>AACGTGCAAT ATTACCAATG TTATTCGTT TATCGTAGTG TTAAAGTGT TTATTTCGTA AAAAAGTGA<br>4061 GCATTCTAGT TGTGGTTTGT CCAAACCTCAT CAATGTATCT TATCATGTCT GTATACCGTC GACCTCTAGC<br>CGTAAGATCA ACACCAACA GGTGAGTA GTTACATAGA ATAGTACAGA CATATGGCAG CTGGAGATCG<br>4131 TAGAGCTGG CGTAATCTG GTCACTAGCTG TTTCCTGTT GAAATTGTTA TCCGCTCACA ATTCCACACA<br>ATCTCGAACC GCATTAGTAC CAGTATCGAC AAAGGACACA CTTAAACAAT AGGGCAGTGT TAAGGTTGT<br>4201 ACATCGAGC CGGAAGCATA AAGTGTAAAG CTTGGGGTC CTAATGAGTC AGCTAACTCA CATTAAATTC<br>TGTATGCTCG GCCTTCGTAAT TTICACATTTC GACCCCCACG GATTACTCAC TCGATTGAGT GTAATTAAACG<br>4271 GTTGGCTCA CTGGCCGGTT TCCAGTCGGG AACCTGTG TGCCAGCTGC ATTAATGAT CGGCCAACGC<br>CAACGGGAGT GACGGGGCAA AGGTAGGCC TTTGGACAGC ACGGTCGAGC TAATTACTTA GCGGGTTGCG<br>4341 GCGGGAGAG GGGGTTGGG TATTGGGGC TCTTCGGCTT CCTCGCTCAC TGACTCGCTG CGCTCGGTG<br>CGCCCTCTC CGCCAAACGC ATAACCCGCG AGAAGGGCAA GGAGGGAGTG ACTGAGCGAC GCGAGCCAGC<br>4411 TTGGGTGCG GCGAGGGTA TCAGCTCACT CAAAGGGGT AATAACGGTTA TCCACAGAAT CAGGGGATAA<br>AAGCCGACGC CGCTCGCCAT AGTCCAGTGA GTTTCGGCCA TTATGCCAAT AGGTGTCITA GTCCCCATT<br>4481 CGCAGGAAAG AACATGTGAG CAAAGGCCA GCAAAAGGCC AGGAACCGTA AAAAGGCCG GTTGTGGCG |

## FIG.9F

|      |             |              |             |              |            |             |              |
|------|-------------|--------------|-------------|--------------|------------|-------------|--------------|
| 4551 | GCGTCCTTTC  | TGTACACTC    | GGTTTCCGGT  | CGTTTCCGG    | TCCTGGCAT  | TTTTCCGG    | CAACGACCGC   |
|      | TTTTTCCATA  | GGCTCCGCC    | CCCTGACGAG  | CATCACAAA    | ATCGACGC   | AAGTCAGAGG  | TGGCGAAACC   |
| 4621 | AAAAGGTAT   | CGAGGGGG     | GGGACTGCTC  | GTAGTGT      | TAGTGTGAG  | TTCACTCTCC  | ACCGCTTTGG   |
| 4691 | CGACAGGACT  | ATAAAAGATAC  | CAGGGGTTTC  | CCCCCTGGAAAG | CTCCCTCGTG | CGCTCTCC    | TTCCGACCCCT  |
|      | GCTGCTCTGA  | TATTCTATG    | GTCCCGCAAAG | GGGGACCTTC   | GAGGAGGAC  | GGCAGAGGAC  | AAGGCTGGGA   |
| 4761 | GCCGCTTAC   | GGATAACCTGT  | CCGCCTTTCT  | CCCCCTGGGA   | AGCGTGGGCC | TTTCTCAATG  | CTCACGCTGT   |
|      | CGGGGAATGG  | CTTATGGACA   | GGCGGAAAGA  | GGGAAGCCCT   | TCGCACCGCG | AAAGAGTTAC  | GAGTGCACAA   |
| 4831 | AGGTATCTCA  | GTTCGGGTGA   | GGTCGTTGCG  | TCCAAGCTGG   | GCTGTGTGCA | CGAACCCCCC  | GTTCAGCCCC   |
|      | TCCATAGAGT  | CAAGCCACAT   | CCAGCAAGCC  | AGGTTCGACCC  | CGACACACGT | GCTTGGGGG   | CAAGTGGGGC   |
| 4901 | AGCAGCCACT  | GGTAACAGGA   | TTAGCAGAGC  | GAGGTATGTA   | GGCGGTGCTA | CAGAGTTCTT  | GAAGTGGTGG   |
|      | TCGTCGGTGA  | CCATTGTCCT   | AATCGTCTCG  | CTCCATACAT   | CCGCACCGAT | GTCTCAAGAA  | CTTCACCAAC   |
| 4971 | CCTAACTACG  | GCTACACTAG   | AAGGACAGTA  | TTTGGGTATCT  | GGGCTCTGCT | GAAGGCCAGTT | ACCTTCGGAA   |
|      | GGATGTGATC  | CGATGTGATC   | TTCCCTGTAT  | AAACCATAGA   | CGCGAGACGA | CTTGGTCAA   | TGGAAGCCTT   |
| 5041 | AAAGAGTTGG  | TAGCTCTTGA   | TCCGGCAAAC  | AAACCAACGC   | TGGTAGGGT  | GGTTTTTTTG  | TTTGCAGCA    |
|      | TTTCTCAACC  | ATCGAGAACT   | AGGCCTTTG   | TTGGTGGCG    | ACCATGCCA  | CCAAAAAAAC  | AAACGTTCTGT  |
| 5111 | GCAGATTAGC  | CGCAGAAAAA   | AAGGATCTCA  | AGAAGATCT    | TTGATCTTT  | CTACGGGGTC  | TGACGCTCAG   |
|      | CGTCATAATGC | GGGTCTTTT    | TTCCCTAGAGT | TCTTCTAGGA   | AACTAGAAAA | GATGCCCGAG  | ACTGCGAGTC   |
| 5181 | TGGAACAAA   | ACTCACGTTA   | AGGGATTTG   | GTCACTGAGAT  | TATCAAAAG  | GATCTTCACC  | TAGATCCTTT   |
|      | ACCTTGCTTT  | TGAGTGCAT    | TCCCCTAAAC  | CAGTACTCTA   | ATAGTTTTTC | CTAGAAAGTGG | ATCTAGGAAA   |
| 5251 | AAAATTAAA   | ATGAAGTTT    | AAATCAATCT  | AAAGTATATA   | TGAGTAAACT | TGGTCTGACA  | GT'LAACAAATG |
|      | ATTTAATTTT  | TACTTCAAA    | TTTGTGTTAGA | TTTCAATATAT  | ACTCATTTGA | ACCGAGCTGT  | CAATGGTTAC   |
| 5321 | CTTATCAGT   | GAGGCACCTA   | TCTCAGCGAT  | CTGTCATTT    | CGTTCATCCA | TAAGTGCCTG  | ACTCCCCGTC   |
|      | GAATTAGTCA  | CCTCCGGTGGAT | AGAGTCGCTA  | GACAGATAAA   | GCAAGTAGGT | ATCAACGGAC  | TGAGGGGCAG   |
| 5391 | GTGTAGATAA  | CTACGATACTG  | GGGGGGCTTA  | CCATCTGGCC   | CCAGTGTGTC | ATGATAACG   | CGAGACCCAC   |
|      | CACATCTATT  | GATGGCTATGC  | CCTCCCGAAT  | GGTAGACCGG   | GGTCACGACG | TIACTATGGC  | GCTCTGGGTG   |
| 5461 | GCTCACCGGC  | TCCAGAGTTA   | TCAGGAAATAA | ACCAAGCCAGC  | CGGAAGGGCC | GAGGCCAGAA  | GTGGCCTGTC   |
|      | CGAGTGGCCG  | AGGTCTAAAT   | AGTCGTTATT  | TGGTCGGTGC   | GCCTTCCGG  | CTCGCGTCTT  | CACCAAGGACG  |
| 5531 | AACTTTATCC  | GGCTCCCATCC  | AGTCATTTAA  | TGTTGGCCGG   | GAAGCTAGAG | TAAGTAGTTC  | GCCAGTTAAT   |
|      | TTGAAATAGG  | CGGAGGTAGG   | TCAGATAATT  | ACAAACGGCC   | CTTGCATCTC | ATTCAATCAAG | CGGTCAATTAA  |

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## FIG.9G

5601 AGTTTGGCA ACAGTTGTGC CATTGCTACA GGCATCGTGG TGTACAGCTC GTTCAGCTC ATGGCTTGTT ATGGCTTCAT  
 TCAAACGGT TGCAACAAACG GTAACGGATGT CGCTAGCACC ACAGTGGAG CAGCAAACCA TACCGAAGTA  
 5671 TCAGCTCCGG TTCCCCAACGA TCAAGGGAG TTAACATGATC CCCCATGTTG TGCAAAAAAG CGGTTAGCTC  
 AGTCGAGGCC AAGGGTTGCT AGTTCGGCTC ATAGTACTAG GGGGTACAAC ACGTTTTTC GCCAATCGAG  
 5741 CTTCGGTCTT CGATCGTGT TCAGAAGTAA GTGGGCCGA GTGTTATCAC TCATGGTTAT GGCAAGCACTG  
 GAAGCCAGGA GGCTAGAAC AGTCTTCATT CAACCGGGGT CACAAATAGTG AGTACCAATA CCGTCGTGAC  
 5811 CATAATTCTC TTACTGTCA GCCATCCGTA AGATGCTTT CTGTAAGCTGG TGAGTACTCA ACCAAGTCA  
 GTATTAGAG ATGACAGTA CGGTAGGCAT TCTACGAAA GACACTGACC ACTCATGAGT TGGTTCAAGTA  
 5881 TCTGAGAATA GTGTATGCCG CGACCGAGTT GCTCTTGCCC GGCCTCAATA CGGGATAATA CCGGCCACAA  
 AGACTCTTACATACGGC GCTGGCTCAA CGAGAACGGG CGCAGTTAT GCCCCTATTAT GGGCGGGTGT  
 5951 TAGCAGAACT TAAAGTGC TCATCATTTGG AAAACGTTCT CGGGGGGAA AACTCTCAAG GATCTTACCG  
 ATCGTCTGAA ATTTCACG AGTAGTAACC TTTTGCAGA AGCCCCGGCTT TTGAGAGTCT CTAGAATGGC  
 6021 CTGTTGAGAT CCAAGTTCGAT GTAAACCCACT CGTGGCACCA ACTGATCTTC AGCATCTTT ACTTTCAACCA  
 GACAAGCTCA GTCAAGCTA CATTGGTGA GCACGTTGGT TGACTAGAAG TCGTAGAAAAA TGAAAGTGGT  
 6091 GCGTTCTGG GTGAGCAAA ACAGGAAGGC AAAATGCCGC AAAAAGGGA ATAAGGGGA CACGGAAATG  
 CGCAAAGACC CACTCGTT TGCCCTTCGG TTTTACGGG TTTTTCCCT TATTCGGCT GTGCCCTTAC  
 6161 TTGAATACTC ATACTCTTC TTTTCAATA TTATTGAAGC ATTATCAGG GTTATTGTTCT CATGAGCGGA  
 AACTTATGAG TATGAGAAGG AAAAGTTAT ATAACACTCG TAAATAGTCC CAATAACAGA GTACTCGCCT  
 6231 TACATATTG ATGTATTAA GAAAATAAA CAATAGGGG TTCCGGCAC ATTTCGGCAC AAAGTGCCAC  
 ATGTATAAAC TACATAAAAT CTTTATTG GTTATCCCCC AAGGGCGGTG TAAAGGGGT TTTCAAGGGT  
 6301 CTGACGTC  
 GACTGCAG